

OIPE

ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/072,622

DATE: 02/27/2002

TIME: 13:50:08

Input Set : A:\07039-331001.TXT

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4 <110> APPLICANT: Chen, Lieping
             Bajorath, Jurgen
      7 <120> TITLE OF INVENTION: ICOS Mutants
     10 <130> FILE REFERENCE: 07039-331001
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/072,622
C--> 12 <141> CURRENT FILING DATE: 2002-02-07
     12 <160> NUMBER OF SEQ ID NOS: 42
     14 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 96
     18 <212> TYPE: PRT
     19 <213> ORGANISM: Mus musculus
     21 <400> SEQUENCE: 1
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                         5
     24 Pro Cys Glu Ser Pro Ser His Asn Thr Asp Val Val Thr Val Leu Gln
                    20
     26 Thr Asn Asp Gln Met Thr Val Ala Thr Thr Phe Thr Glu Lys Asn Thr
     27
                                    40
     28 Val Gly Phe Leu Asp Tyr Pro Phe Ser Gly Thr Phe Asn Glu Ser Arg
     30 Val Asn Leu Thr Ile Gln Gly Leu Arg Ala Val Asp Gly Leu Tyr Leu
                            70
     32 Cys Val Leu Phe Val Gly Met Gly Gln Ile Tyr Val Ile Pro Glu Pro
     35 <210> SEQ ID NO: 2
     36 <211> LENGTH: 96
     37 <212> TYPE: PRT
     38 <213> ORGANISM: Rattus norvegicus
     40 <400> SEQUENCE: 2
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     42 1
                         5
     43 Pro Cys Glu Ala Ser Ser His Asn Thr Asp Val Val Thr Val Leu Gln
                                         25
     45 Thr Asn Asp Gln Val Thr Val Ala Thr Thr Phe Thr Val Lys Asn Thr
     47 Leu Gly Phe Leu Asp Asp Pro Phe Ser Gly Thr Phe Asn Glu Ser Arg
                                55
     49 Val Asn Leu Thr Ile Gln Gly Leu Arg Ala Ala Asp Gly Leu Tyr Phe
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     51 Cys Val Leu Phe Val Gly Met Gly Gln Ile Tyr Val Ile Pro Glu Pro
     54 <210> SEQ ID NO: 3
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55 <211> LENGTH: 96
56 <212> TYPE: PRT
57 <213> ORGANISM: Homo sapiens
59 <400> SEQUENCE: 3
60 Val Ala Gln Pro Ala Val Leu Ala Ser Ser Arg Gly Ile Ala Ser Phe
62 Val Cys Glu Ala Ser Pro Gly Lys Ala Thr Val Val Thr Val Leu Gln
64 Ala Asp Ser Gln Val Thr Val Ala Ala Thr Tyr Met Met Gly Asn Glu
                               40
65 35
66 Leu Thr Phe Leu Asp Asp Ser Ile Thr Gly Thr Ser Ser Gly Asn Gln
                          - 55
68 Val Asn Leu Thr Ile Gln Gly Leu Arg Ala Met Asp Gly Leu Tyr Ile
70 Cys Val Leu Tyr Leu Gly Ile Gly Gln Ile Tyr Val Ile Pro Glu Pro
73 <210> SEQ ID NO: 4
74 <211> LENGTH: 96
75 <212> TYPE: PRT
76 <213> ORGANISM: Bos taurus
78 <400> SEQUENCE: 4
79 Val Ser Gln Pro Ala Val Leu Ala Ser Ser Arg Gly Val Ala Ser Phe
81 Val Cys Glu Ala Ser Ser His Lys Ala Thr Val Val Thr Val Leu Gln
83 Ala Asn Ser Gln Met Thr Val Ala Met Thr Tyr Thr Val Glu Asn Glu
                               40
85 Leu Thr Phe Ile Asp Asp Ser Thr Thr Gly Ile Ser His Gly Asn Lys
                           55
87 Val Asn Leu Thr Ile Gln Gly Leu Ser Ala Met Asp Gly Leu Tyr Ile
89 Cys Val Leu Tyr Val Gly Met Gly Gln Ile Tyr Val Ile Pro Glu Pro
92 <210> SEQ ID NO: 5
93 <211> LENGTH: 95
94 <212> TYPE: PRT
95 <213> ORGANISM: Mus musculus
97 <400> SEQUENCE: 5
98 Val Gln Pro Leu Val Asp Ser Glu Ser Leu Cys Leu Ala Ala Leu Tyr
100 Gly Val Asn Asp Val Gly Asn Gly Asn Phe Thr Tyr Gln Pro Gln Phe
102 Arg Ser Asn Ala Glu Phe Asn Cys Asp Gly Asp Phe Asp Asn Glu Thr
                                40
104 Val Thr Phe Arg Leu Trp Asn Leu His Val Asn His Thr Asp Ile Tyr
106 Phe Cys Lys Ile Glu Phe Met Tyr Pro Pro Pro Tyr Leu Asp Asn Glu
108 Arg Ser Asn Gly Thr Ile Ile His Ile Lys Glu Lys His Leu Cys
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111	<210> SI	EQ ID	NO:	. 6												
112	<211> L1	ENGTH	: 95	5												
113	<212> TYPE: PRT															
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116	<400> SI	EQUEN	CE:													
	Val Gln 1				Asp	Asn	Glu	Ser	Leu 10	Cys	Leu	Ala	Ala	Leu 15	Tyr	
	Gly Val		Asp 20	Val	Gly	Asn	Gly	Asn 25	Phe	Thr	Tyr	Gln	Pro 30	Gln	Phe	
	Arg Pro		-	Gly	Phe	Asn	Cys		Gly	Asn	Phe	Asp		Glu	Thr	
122		35					40					45				
123 124	Val Thr 50	Phe	Arg	Leu	Trp	Asn 55	Leu	Asp	Val	Asn	His 60	Thr	Asp	Ile	Tyr	
	Phe Cys	Lvs	Ile	Glu	Val	Met	Tyr	Pro	Pro	Pro	Tyr	Leu	Asp	Asn	Glu	
126	_	4			70		-			75	-		-		80	
	Lys Ser	Asn	Glv	Thr	Ile	Ile	His	Ile	Lvs	Glu	Lys	His	Leu	Cys		
128	270 002		u = 1	85					90		-1-			95		
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	<211> LI															
	<212> TY															
	<213> OI			Bos	tauı	rus										
	<400> SI															
	Val Gln				Asn	Asn	Glu	Asn	Leu	Cvs	Phe	Ser	Ala	Leu	Tvr	
137				5			0 _u		10	-1-				15	-1-	
	Gly Ala	Asp	Ala	_	Val	Asn	Glv	Asn		Ser	His	Pro	His		Phe	
139	0-1 11-0	_	20				1	25					30			
	His Ser			Glv	Phe	Asn	Cvs		Gly	Lys	Leu	Gly	Asn	Glu	Thr	
141		35		1			40		1			45				
	Val Thr	Phe	Tvr	Leu	Lvs	Asn	Leu	Tyr	Val	Asn	Gln	Thr	Asp	Ile	Tyr	
143	50		- 4		-	55		1			60		-		-	
144	Phe Cys	Lys	Ile	Glu	Val	Met	Tyr	Pro	Pro	Pro	Tyr	Leu	Asp	Asn	Glu	
145		-			70		-			75	-		_		80	
146	Lys Ser	Asn	Gly	Thr	Ile	Ile	His	Val	Lys	Glu	Gln	His	Phe	Cys		
147	_		_	85					90					95		
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151	<212> TY	YPE:	PRT													
152	<213> OF	<pre><213> ORGANISM: Homo sapiens</pre>														
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	Val Gln				Tyr	Asp	Ala	Asn	Leu	Cys	Phe	Ser	Ala	Leu	His	
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	Gly Leu	Asp .	Ala	Val	Val	Tyr	Gly	Asn	Tyr	Ser	Gln	Gln	Leu	Gln	Val	
158			20			•	- 4	25	-				30			
	Tyr Ser			Glv	Phe	Asn	Cys		Gly	Lys	Leu	Gly		Glu	Ser	
160	<u>.</u>	35		4			40	-	_	-		45				
	Val Thr		Tyr	Leu	Gln	Asn	Leu	Tyr	Val	Asn	Gln		Asp	Ile	Tyr	
162	50					55		4 -			60		•		-	
	Phe Cys	Lys	Ile	Glu	Val		Tyr	Pro	Pro	Pro	Tyr	Leu	Asp	Asn	Glu	
	- 2 -						-				-		•			

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80
                        70
165 Lys Ser Asn Gly Thr Ile Ile His Val Lys Glu Lys His Leu Cys
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168 <210> SEQ ID NO: 9
169 <211> LENGTH: 112
170 <212> TYPE: PRT
171 <213> ORGANISM: Mus musculus
173 <400> SEQUENCE: 9
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176 Cys Lys Tyr Pro Asp Ile Val Gln Gln Leu Lys Met Arg Leu Phe Arg
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178 Glu Arg Glu Val Leu Cys Glu Leu Thr Lys Thr Lys Gly Ser Gly Asn
                                40
180 Ala Val Ser Ile Lys Asn Pro Met Leu Cys Leu Tyr His Leu Ser Asn
                            55
182 Asn Ser Val Ser Phe Phe Leu Asn Asn Pro Asp Ser Ser Gln Gly Ser
                                            75
                        70
183 65
184 Tyr Tyr Phe Cys Ser Leu Ser Ile Phe Asp Pro Pro Pro Phe Gln Glu
                                        90
                    85
186 Arg Asn Leu Ser Gly Gly Tyr Leu His Ile Tyr Glu Ser Gln Leu Cys
                                    105
187
189 <210> SEQ ID NO: 10
190 <211> LENGTH: 111
191 <212> TYPE: PRT
192 <213> ORGANISM: Homo sapiens
194 <400> SEQUENCE: 10
195 Ala Asn Tyr Glu Met Phe Ile Phe His Asn Gly Gly Val Gln Ile Leu
197 Cys Lys Tyr Pro Asp Ile Val Gln Gln Phe Lys Met Gln Leu Leu Lys
                                     25
199 Gly Gly Gln Ile Leu Cys Asp Leu Thr Lys Thr Lys Gly Ser Gly Asn
                                40
201 Thr Val Ser Ile Lys Ser Leu Lys Phe Cys His Ser Gln Leu Ser Asn
203 Asn Ser Val Ser Phe Phe Leu Tyr Asn Leu Asp His Ser His Ala Asn
                        70
205 Tyr Tyr Phe Cys Asn Leu Ser Ile Phe Asp Pro Pro Pro Phe Lys Val
                    85
                                        90
207 Thr Leu Thr Gly Gly Tyr Leu His Ile Tyr Glu Ser Gln Leu Cys
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210 <210> SEQ ID NO: 11
211 <211> LENGTH: 6
212 <212> TYPE: PRT
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 11
216 Met Tyr Pro Pro Pro Tyr
217 1
219 <210> SEQ ID NO: 12
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Input Set : A:\07039-331001.TXT

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220 <211> LENGTH: 199
221 <212> TYPE: PRT
222 <213> ORGANISM: Homo sapiens
224 <400> SEQUENCE: 12
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227 Val Leu Thr Gly Glu Ile Asn Gly Ser Ala Asn Tyr Glu Met Phe Ile
                20
229 Phe His Asn Gly Gly Val Gln Ile Leu Cys Lys Tyr Pro Asp Ile Val
231 Gln Gln Phe Lys Met Gln Leu Leu Lys Gly Gln Ile Leu Cys Asp
233 Leu Thr Lys Thr Lys Gly Ser Gly Asn Thr Val Ser Ile Lys Ser Leu
                                            75
                        70
235 Lys Phe Cys His Ser Gln Leu Ser Asn Asn Ser Val Ser Phe Phe Leu
                                        90
                    85
237 Tyr Asn Leu Asp His Ser His Ala Asn Tyr Tyr Phe Cys Asn Leu Ser
                                    105
                100
239 Ile Phe Asp Pro Pro Pro Phe Lys Val Thr Leu Thr Gly Gly Tyr Leu
                                120
                                                     125
           115
241 His Ile Tyr Glu Ser Gln Leu Cys Cys Gln Leu Lys Phe Trp Leu Pro
                                                 140
                            135
242
243 Ile Gly Cys Ala Ala Phe Val Val Val Cys Ile Leu Gly Cys Ile Leu
                                             155
                        150
245 Ile Cys Trp Leu Thr Lys Lys Lys Tyr Ser Ser Ser Val His Asp Pro
                                         170
247 Asn Gly Glu Tyr Met Phe Met Arg Ala Val Asn Thr Ala Lys Lys Ser
                                     185
                180
249 Arg Leu Thr Asp Val Thr Leu
            195
250
252 <210> SEQ ID NO: 13
253 <211> LENGTH: 62
254 <212> TYPE: DNA
255 <213> ORGANISM: Artificial Sequence
257 <220> FEATURE:
258 <223> OTHER INFORMATION: Primer
260 <400> SEQUENCE: 13
261 ttatgcaaat atcctgacat tgtcagggag ttccggatgc agttgctgaa aggggggcaa
                                                                             60
                                                                             62
262 at
264 <210> SEQ ID NO: 14
265 <211> LENGTH: 22
266 <212> TYPE: DNA
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Primer
272 <400> SEQUENCE: 14
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273 gacaatgtca ggatatttgc at
275 <210> SEQ ID NO: 15
276 <211> LENGTH: 59
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VERIFICATION SUMMARY

DATE: 02/27/2002

PATENT APPLICATION: US/10/072,622

TIME: 13:50:09

Input Set : A:\07039-331001.TXT
Output Set: N:\CRF3\02272002\J072622.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date